

APPROVED	O. G. FIG.
BY	CLASS SUBCLASS
DRAFTSMAN	

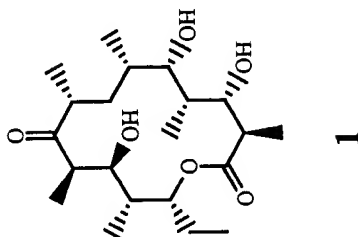
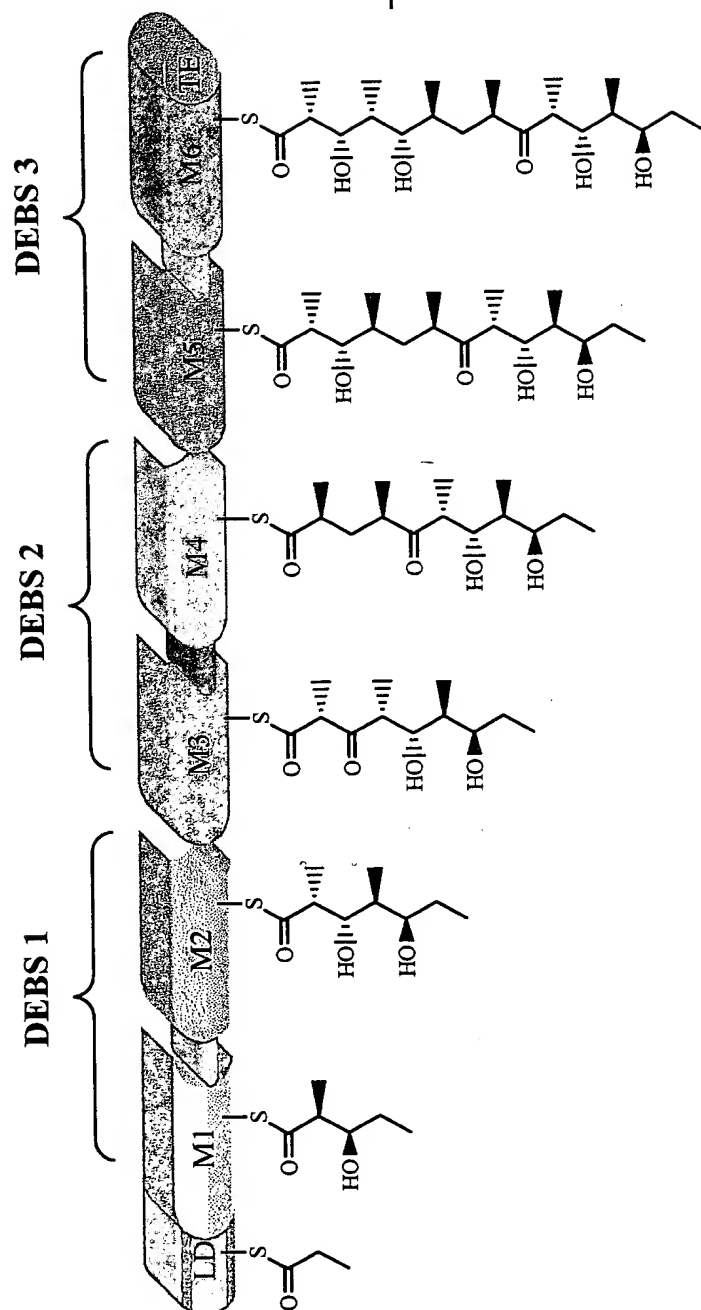


Figure 1

APPROVED	O.G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

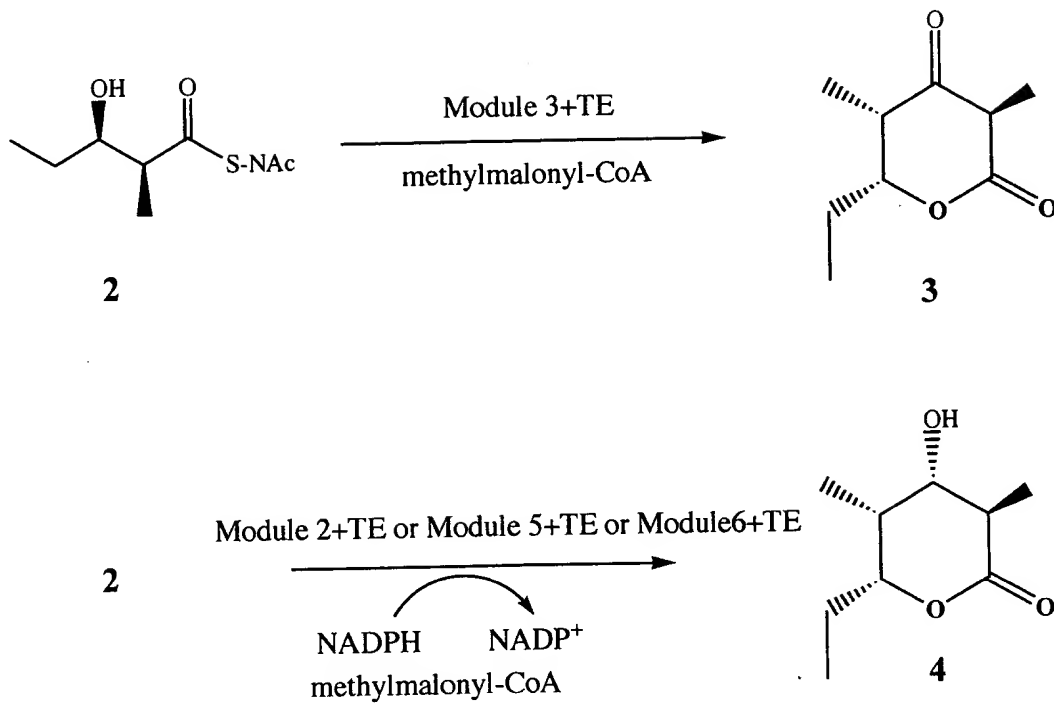


Figure 2

APPROVED	O G. FIG.	
BY	CLASS	SUBCLASS
DRAFTSMAN		

(a) INTRA-POLYPEPTIDE LINKER

M2ery: GGATGAEQAAPATT..APVD
M4ery: VGDAD..QAA.VRVVGAA.DES
M6ery: VGAAEAEQA.PALVREVPKAD
M2rif: FGSA.A.NR.PAEIGTAAAE
M3rif: LG..ER.PAAPAPVTRDVSD
M5rif: GETVAGAPATPVTTVADAG
M3rap: .ELFTGENPAPVRGPVSAVGQD
M4rap: .ELFTGENPAPVRGPVSVVGQD
M7rap: .ELFTGENPAPVRGPVSA.GQD

(b) N-TERMINAL INTER-POLYPEPTIDE LINKER

M3ery:VTD SE KVAEYLRR .ATL DLRAAR QRIRE..LES
M5ery: MSGDNGM.TE E.KLRRYLKR TVT.ELDSVT ARLRE..VEH RAG
M4rif:MSAPNE QIVDAL.R ASLKE....N VRLOQENSAL AAAAA
M7rif:VSASYE KVVEAL.R KSLEE....V GTLKKRNRQL ADAAG
M8rif:V.AD EGQLRDYLKR .AIADARDAR TRLRE..VEE QAR
M9rif:MATD E.KLLKYLKR .VTAE LHS... ..LRKQGARH .AD
M5rap:MR.. EDQLLDAL.R KSVKE....N ARLRKANTSL RAAMD
M11rap:M.PEQD KVVEYL.R WATAELHTTR AKL.....EA LAAANT

(a) INTRA-POLYPEPTIDE LINKER

M2ery: GGATGAEQAAPATT..APVD (SEQ ID NO:3)
M4ery: VGDAD..QAA.VRVVGAA.DES (SEQ ID NO:4)
M6ery: VGAAEAEQA.PALVREVPKAD (SEQ ID NO:5)
M2rif: FGSA.A.NR.PAEIGTAAAE (SEQ ID NO:6)
M3rif: LG..ER.PAAPAPVTRDVSD (SEQ ID NO:7)
M5rif: GETVAGAPATPVTTVADAG (SEQ ID NO:8)
M3rap: .ELFTGENPAPVRGPVSAVGQD (SEQ ID NO:9)
M4rap: .ELFTGENPAPVRGPVSVVGQD (SEQ ID NO:10)
M7rap: .ELFTGENPAPVRGPVSA.GQD (SEQ ID NO:11)

(b) N-TERMINAL INTER-POLYPEPTIDE LINKER

M3ery:VTD SE KVAEYLRR .ATLIDLRAAR QRIRE..LES (SEQ ID NO:12)
M5ery: MSGDNGM.TE E.KLRRYLKR TVT.ELDSVT ARLRE..VEH RAG (SEQ ID NO:13)
M4rif:MSAPNE QIVDAL.R ASLKE....N VRLQQENSAL AAAAA (SEQ ID NO:14)
M7rif:VSASYE KVVEAL.R KSLEE....V GTLKKRNRQL ADAAG (SEQ ID NO:15)
M8rif:V.AD EGQLRDYLKR .AIADARDAR TRLRE..VEE QAR (SEQ ID NO:16)
M9rif:MATD E.KLLKYLKR .VTAELES.. ..LRKQGARE .AD (SEQ ID NO:17)
M5rap:MR.. EDQLLDAL.R KSVKE....N ARLRKANTSL RAAMD (SEQ ID NO:18)
M11rap:M.PEQD KVVEYL.R WATAELHTTR AKL.....EA LAAANT (SEQ ID NO:19)

Figure 3